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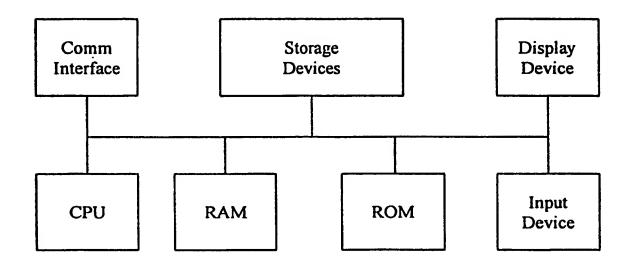
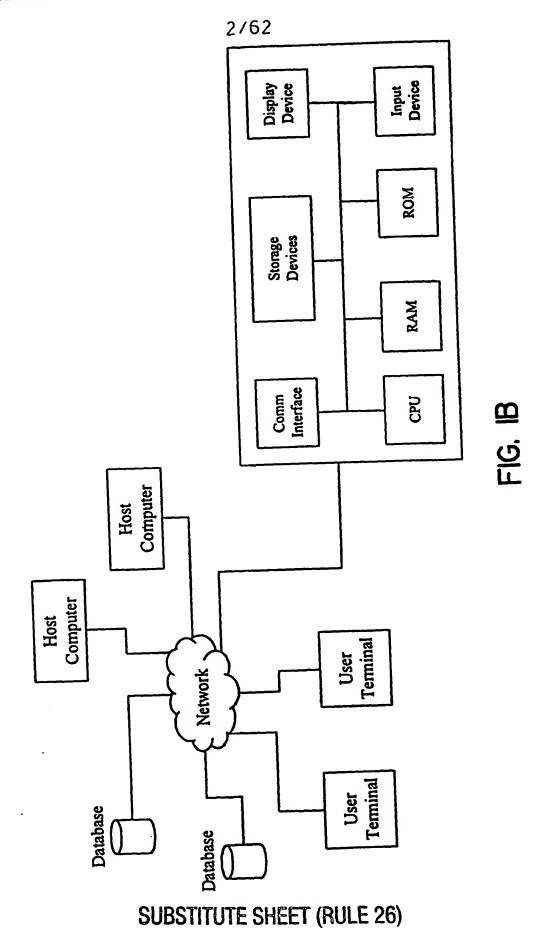
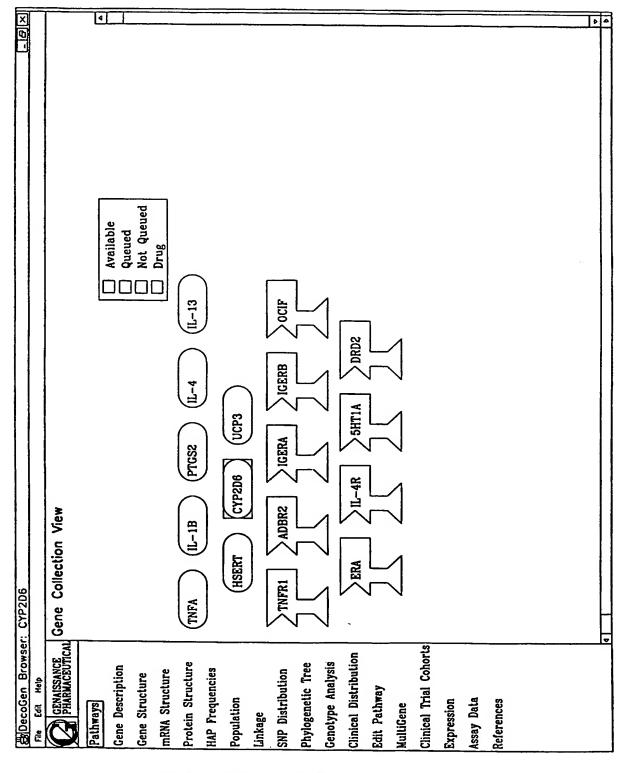


FIG. IA

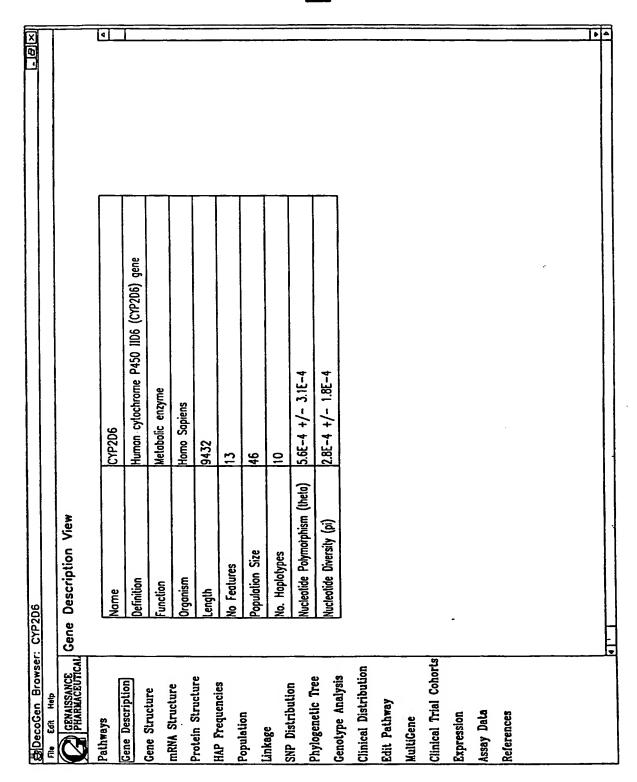


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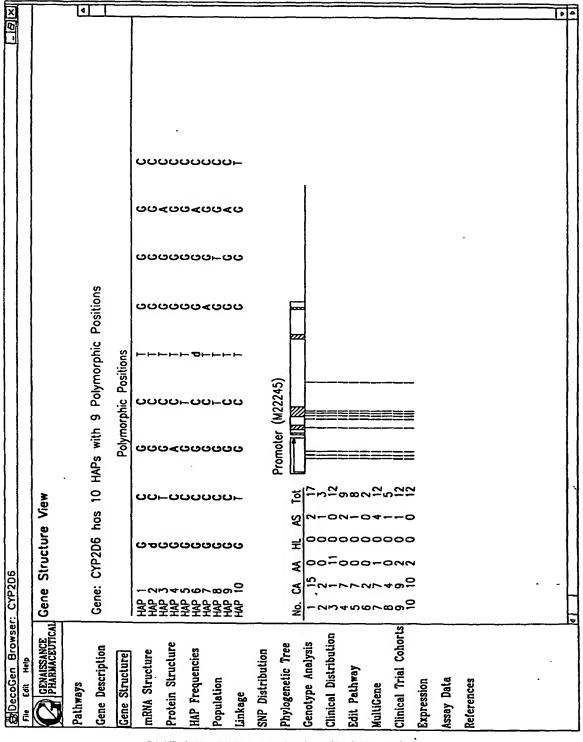
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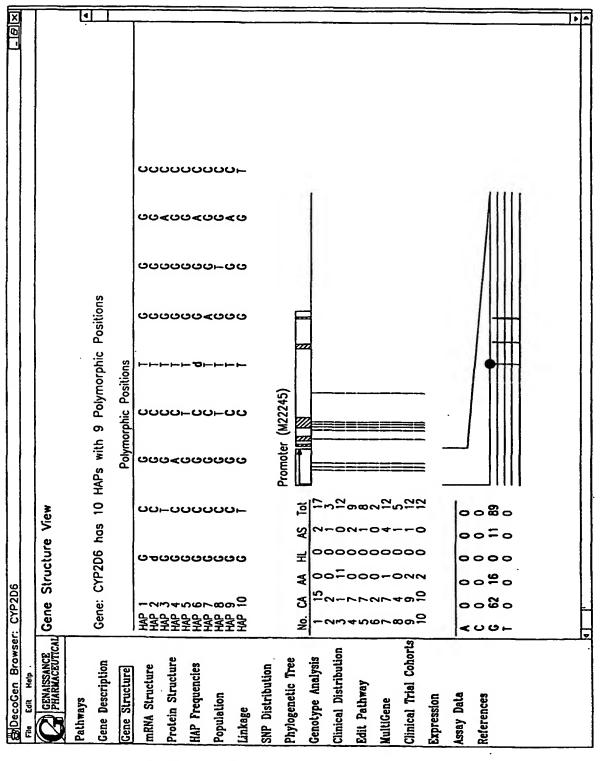


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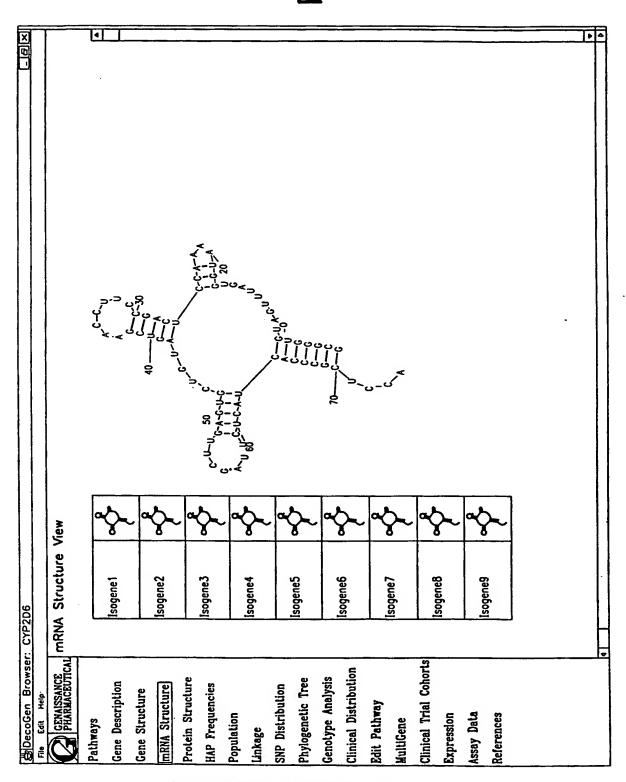
FIG. 4b .25



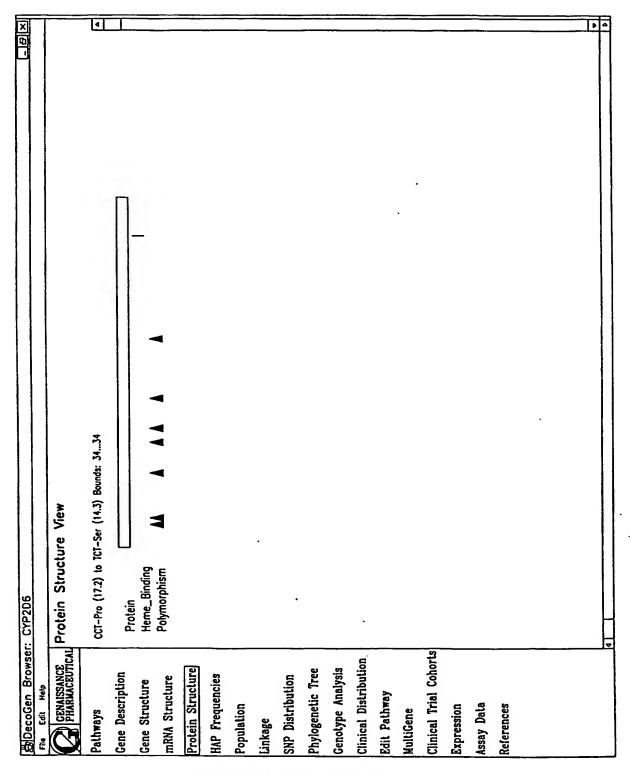
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GENAISSANCE PHARMACEUTICAL	Population View	ew.						
Pathways	PIO	Ethnicity	Age	Gender	HAP 1	HAP 2	Test .	<u>II</u>
Gene Description	UP070	CA	66	L.	GCGCTGGGC	GCGCTGGGC	0.1	
Gene Structure	UP072	CA.	66	LL.	dcccTcccc	GCACTGGGC	0.2	
mRNA Structure	UP074	æ	66	٤	GCGCTGGGC	GCACTGGGC	0.2	
Protein Structure	UP132	CA	66	×	GTGCTGGGT	GTGCTGGGT	0.3	
HAP Frequencies	UP133	క	66	×	GCGCTGGGC	GTGCTGGGT	0.2	
Population	UP134	S	66	4	GCGCTGGGC	cccrcccc	0.1	
Linkage	UP137	క	66	≖	GTGCTGGGT	GTGCTGGGT	0.1	
SNP Distribution	UP009	క	66	LL.	GCACTGGGC	GCACTGGGC	0.1	
Phylogenetic Tree	UP014	క	66	Ŀ	decerecee	GCGCTGGAC	0.3	
Genotype Analysis	UP020	క	66	L.	GTGCTGGGT	GTGCTGGGT	0.2	
Clinical Distribution	UP021	ర	66	3	СССТСТСС	GCGCTGGGC	0.4	
Edit Pathway	UP022	হ	66	2	GCGCTGGAC	GCGCTGGAC	0.3	
MultiGene	OR001	AS	99	æ	desertesses	ccincrec	0.2	
Clinical Trial Cohorts	OR002	AS	66	×	GCGCTGGGC	GCGCTGGC	0.3	
Expression	0R004	AS	66	<u>u</u>	GCACTGGGC	GCACTGGGC	0.5	
Assay Data	08006	S	66		ссеттесес	GCGCTGGAC	0.1	
References	WT003	ర	66	<u>ب</u>	ссетсесс	СССТСТССС	0.2	
•	WT005	ర	66	×	ссеттесес	GCGCTGGGC	0.2	
	WT007	ర	66	≆	ссепетес	ссепстос	0.4	
	UP012	ర	66	<u>ı.</u>	GCGCTAGGC	GCCTGCAC	0.1	
	UP135	ర	66	₹	GCACTGGGC	GCCTCCAC	0.2	ŀ
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